

**AMENDMENTS TO THE DRAWINGS:**

The attached sheet of drawings includes a correction to Figure 1. The corrected drawing places the number 40 above the amino acid located at this position. In original Figure 1, the number 40 was placed above the amino acid located at position 39.

Attachment: Replacement Sheet  
Annotated Sheet Showing Changes

Application No. 09/976,414  
Amendment Dated September 16, 2003  
Replacement Sheet

No: 1            10            20            30            40            50  
a) AQSVPYGVSQIKAPALHSQGYTGSNVKVAVIDSGIDSSHPDLKVAGGASM  
b) AQSVPWGISRVQAPAAHNRGLTGSGVKVAVLDTGI\*STHPDLNIRGGASF

No:            60            70            80            90            100  
a) VPSETNPFQDNNSHGTHVAGTVAALNNSIGVLGVAPSASLYAVKVLGADG  
b) VPGEPT\*QDGNHGHGTHVAGTVAALNNSIGVIGVAPSADLYAVKVLGANG

No:            110            120            130            140            150  
a) SGQYSWIINGIEWAIANNMDVINMSLGGPSGSAALKAAVDKAVASGVVVV  
b) RGSVSGIAQGLEWAAANNMHIANMSLGSDAPSTTLERAVNYATSQGV LVI

No:            160            170            180            190            200  
a) AAAGNEGTSGSSSTVGYPGKYPSVIAVGAVDSSNQRASFSSVGPELDVMA  
b) AATGNNG\*SGS\*\*\*VGYPARYANAMAVGATDQNNRRANFSQYGTGIDIVA

No:            210            220            230            240            250  
a) PGVSIQSTLPGNKYGAYNGTSMASPHVAGAAALILSKHPNWTNTQVRSSL  
b) PGVDIESTYPGSSYDSLSTMATPHVAGVAALVKQKNPSWSNVQIRNHL

No:            260            270            275  
a) ENTTTKLGDSFYYGKGLINVQAAAQ  
b) KNTATSLGSTNLYGSGLVNAEAATR

Fig. 1

Application No. 09/976,414  
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No: 1	10	20	30	40	50
a) AQSVPYGVSQIKAPALHSQGYTGSNVKVAVIDSGIDSSHPLKLVAGGASM					
b) AQSVPWGISRVQAPAAHNRGLTGSGVKVAVLDTGI*STHPDLNIRGGASF					
No:	60	70	80	90	100
a) VPSETNPFQDNNSHGTHVAGTVAALNNSIGVLGVAPSASLYAVKVLGADG					
b) VPGEPT*QDGNHGTHVAGTVAALNNSIGVIGVAPSADLYAVKVLGANG					
No:	110	120	130	140	150
a) SQQYSWIINGIEWAIANNMDVINMSLGGPSGSAALKA AVDKAVASGVVVV					
b) RGSVSGIAQGLEWAAANNMHIANMSLGSDAPSTTLERAVNYATSQGV LVI					
No:	160	170	180	190	200
a) AAAGNEGTS GSSSTVGYPGKYPSVIAVGAVDSSNQ RASFSSVGP ELDVMA					
b) AATGNNG*SGS***VGYPARYANAMAVGATDQNNRRANFSQYGTGIDIVA					
No:	210	220	230	240	250
a) PGVSIQSTLPGNKYGAYNGTSMASPHVAGAAALILSKHPNWTNTQVRSSL					
b) PGVDIESTYPGSSYDSLSTMATPHVAGVAALVKQKNPSWSNVQIRNHL					
No:	260	270	275		
a) ENTTTKLGDSFYYGKGLINVQAAAQ					
b) KNTATSLGSTNLYGSLVNAEAAATR					

Fig. 1